Detection of Human Metapneumovirus in Hospitalized Children with Acute Respiratory Tract Infections in Sulaimani Province, Iraq

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Abstract

**Background:** Human Metapneumovirus (hMPV) is a member of Paramyxoviridae family. It is an important viral pathogen of respiratory tract infections among children. Therefore, the aim of this study is to determine the incidence of hMPV infections among children less than 5 years of age with respiratory tract infections at the pediatric hospital in Sulaimani city of Iraq. Also, this study aims to evaluate the different diagnostic methods for the detection of this virus.

**Method:** Nasopharyngeal swabs and throat swabs were collected from 300 hospitalized children with respiratory infections under 5 years of age between April 2011 and March 2012. Each sample was used for hMPV detection by conventional reverse transcriptase (RT-PCR) and direct fluorescent assay (DFA). A questionnaire which was designed for acute respiratory tract infections among hospitalized children was also included.

**Results:** Human Metapneumovirus was detected in 16% by RT-PCR and 14.7% by DFA. Autumn – winter period was the most common season for hMPV infections with its peak occurrence in January and February.

**Conclusions:** hMPV is an important pathogen associated with RTIs in children. RT-PCR is highly sensitive and specific for the detection of hMPV than antigen detection methods used for the diagnosis of hMPV viruses.

**Keywords:** Human metapneumovirus; Hospitalized children; Direct fluorescent assay; RT-PCR

Introduction

Human Metapneumovirus (hMPV) is an enveloped single stranded negative-sense RNA virus [1]. It is a member of the Paramyxoviridae family, subfamily Pneumovirinae, and genus Metapneumovirus [2]. The hMPV was first identified in Netherland by Van den Hoogen, et al. However, the virus was first isolated from stored nasopharyngeal aspirates of infected children collected over a 20 year period [3].

hMPV appears to have respiratory epithelium tropism [4]. The symptoms of both upper and lower respiratory tract disease have been associated with hMPV infections in infants, young children, elderly, and immunocompromized patients. However, at the age of five years, virtually every individual has experienced at least one hMPV infection [5-7]. In young hospitalized children, the clinical features associated with hMPV infections are similar to those of RSV [8].

There are mainly two major techniques for the detection of hMPV from respiratory specimens, conventional reverse transcriptase – PCR (RT-PCR) and immunofluorescence technique, although shell vial culture and routine tissue culture can also be use [9,10].

The laboratory diagnostic methods that are routinely used in our hospitals do not usually include the diagnosis of hMPV in their schedule investigations. Therefore, to the best of our knowledge, data on the incidence and clinical importance of hMPV infections in children with ARTI in Sulaimani province of Iraq is limited.

This study aims to determine the epidemiological characteristics of hMPV infections among hospitalized children with ARTI and immunocompromized children from Hiwa hospital in Sulaimani province of Iraq.

Materials and Methods

Study design

Cross sectional, hospital – based study.

Keywords: Human metapneumovirus; Hospitalized children; Direct fluorescent assay; RT-PCR

Introduction

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This study aims to determine the epidemiological characteristics of hMPV infections among hospitalized children with ARTI and immunocompromized children from Hiwa hospital in Sulaimani province of Iraq.

Patients

Three hundred hospitalized children were involved in this study: two hundred and seventy hospitalized children less than five years of age, were admitted to Pediatric Teaching Hospital in Sulaimani city for respiratory tract infections of undiagnosed etiology; while thirty children were immunocompromized, suffering from RTIs in Hiwa Hospital. Both genders were included and were chosen by random sampling. The specimen collection began from April 2011 to March 2012. In addition, a signed informed consent was obtained from the legal guardians of the children who participated in this study.

Questionnaire

The patients were subjected to a questionnaire that included demographical and clinical data related to respiratory infections.

Specimens

Two Nasopharyngeal swabs (NPSs) were taken from each child by flocked swab applicators (Micro Rheologics Company/Italy). One NPS was placed immediately after collection in 3 ml viral transport medium (VTM), and then VTM was stored at -70°C. The other swab was immediately rolled on the labeled slide. In addition to NPSs, two throat swabs (TSs) were also collected from each child, and these TSs were manipulated in the same manner as NPSs.

Methods

Both direct immunofluorescence assay (DFA) and conventional
reverse transcriptase–PCR (RT–PCR) were used for the detection of hMPV from the swabs collected from the children in this study.

**DFA**

DFA was performed according to the manufacturer’s instructions (Light Diagnostics, Millipore, USA). In summary, the nasopharyngeal swab was rolled on the slide, and the epithelial cells were fixed in cold 100% acetone for 10 minutes at 2-8°C. The spot of specimen on slide was bordered by a marker pen. Thus, about 40 µl of SIMULFLUOR HMPV/RSV reagent (conjugate) was used to cover the spot of the smear. The slide was incubated for 15 minutes at 37°C in a humid chamber, and then it was rinsed thoroughly with PBS/Tween 20 solution, and allowed to air dry. The slide was then mounted under a cover slip using an aqueous mounting medium. Microscopical examination of slide was performed using a fluorescence microscope (Olympus BX60; Olympus Optical Co. Ltd., Tokyo, Japan) with 100-400x magnification for cells exhibiting the apple green fluorescence of fluorescein isothiocyanate (FITC) stain for hMPV positive slides.

**Molecular technique**

The conventional reverse transcriptase–PCR (RT–PCR) was the molecular technique for the detection of hMPV. The technique targeted the F gene of hMPV genome using primers designed in this study (Table 1). However, it was taken from NCBI Reference Sequence: NC_004148.2.

Therefore, the technique was performed in the following sequence:

1. Extraction of viral RNA form NPS using viral DNA/RNA extraction kit (Intron Biotechnology Company/South Korea), and was performed according to the manufacturer's instruction.

2. Reverse transcription was done with two–steps reverse transcriptase–PCR kit in thermal cycler (TC312/ Techne Company/United Kingdom) for the synthesis of first strand cDNA; and it was performed according to the manufacturer's instruction.

3. PCR amplification: The PCR amplification protocol which is mentioned in Table 2, was used in this study. Also, ready to use positive control was used to check the third step.

4. Horizontal gel electrophoresis (Phero – Sub 1010-E/ Biotec – Fischer Company/USA) was used for the detection of PCR products and the amplicons were verified using DNA ladder, and then, the amplicons were viewed in Gel documentation (Uvido/ Uvitec Company/UK). The components were added in one micro centrifuge tube for a total of 25 µl reaction volume. Positive control is added in a separate tube with its primers.

**Table 1:** Primers used in this study.

<table>
<thead>
<tr>
<th>Gene</th>
<th>Polarity of primer</th>
<th>Nucleotide position of primer</th>
<th>Primer sequence (5’→3’)</th>
<th>Amplicon size</th>
</tr>
</thead>
<tbody>
<tr>
<td>hMPV F gene</td>
<td>F</td>
<td>3796-3815</td>
<td>ATGTTGGAGAACCGTGCGAT</td>
<td>465</td>
</tr>
<tr>
<td></td>
<td>R</td>
<td>4260-4241</td>
<td>CCCCCTCTCTGTTGCTGCCAA</td>
<td></td>
</tr>
</tbody>
</table>

**Results**

The DFA technique for the detection of hMPV antigen in epithelial cells of nasopharynx showed that positive specimens were emitting green fluorescence as they were stained with fluorescein isothiocyanate (FITC) dye, and were examined under a fluorescent microscope (Figures 1 and 2). The results revealed that 44 (14.7%) NPS specimens were hMPV positive, while the remaining 256 (85.3%) patients were negative. Also, the results of the TS specimens revealed 41 (13.7%) positive and 259 (86.7%) negative cases. Therefore, the DFA results showed that the differences between NPS and TS were not statistically significant (P=0.8150) (Figure 3).

Conventional RT-PCR was performed for the detection of hMPV from the collected specimens. The results revealed that 48 (16%) patients were positive, while the remaining 252 (84%) were negative.

**Table 2:** The protocol for PCR amplification.

<table>
<thead>
<tr>
<th>Component</th>
<th>Volume</th>
</tr>
</thead>
<tbody>
<tr>
<td>Master mix</td>
<td>12.5 µl</td>
</tr>
<tr>
<td>Forward primers</td>
<td>3 µl</td>
</tr>
<tr>
<td>Reverse primers</td>
<td>3 µl</td>
</tr>
<tr>
<td>CDNA</td>
<td>6.5 µl</td>
</tr>
</tbody>
</table>
Thus, each reaction tube for each patient represents a combination of NPS and TS specimens.

All hospitalized immunocompromized patients were tested for hMPV infections, and the results showed that 4 (13.3%) patients from the total number of 30 patients were positive, while the remaining 26 (86.7%) were negative for hMPV as tested by RT-PCR. The same results were also obtained when specimens were tested by DFA. Hence, the differences in results between the immunocompromized and immunocompetent patients are not statistically significant (p=0.7983) (Table 3).

The total number of hospitalized children that participated in this study was 300, and all of them were under 5 years of age. 48 (16%) were positive for hMPV when tested by RT-PCR, and 252 (84%) were negative. The 48 hMPV positive patients were mostly males and were 34 (70%), whereas 14 (29.2%) of them were females. Hence, the male to female ratio was 2.4:1.

All children were divided into five groups, <1, 1 to < 2, 2 to <3, 3 to <4, and 4 to <5. The highest hMPV infection detected by RT-PCR was in the age group <1 years old, while the lowest frequency was in 4 to <5.

Abnormal breathing sounds 40 120 <0.0001
Added breathing sounds 28 88 0.0123
Cyanosis 3 33 0.2300
History of chronic disease 4 31 0.6235
History of previous similar condition 23 125 0.2650
History of contact with similar condition 16 31 0.0030
Family history of chronic respiratory disease 3 21 0.2698
Mean hospital stay (days) 4.0 3.3 0.1168
BMI
Normal 41 213
Above normal 1 21 0.1654
Below normal 6 18

WBCs count
High total 42 165 0.0020
Granulocytosis 29 108 0.7178
Lymphocytosis 10 48 0.5674

Clinical illness
Bronchiolitis 13 66 0.0078
Pneumonia 21 60
Others 14 126

SpO₂ (%)
Normal 8 130 <0.0001
Subnormal 40 114

Chest x-ray
Normal 14 145 0.0004
Abnormal 34 107

Type of treatment
Supportive 48 252
Specific antibacterial drugs 48 250
Specific antiviral drugs 0 0

Condition at discharge
Complete recovery 3 19 1.000
Discharged with medication 45 230 1.000
Death 0 3

Table 4: Demographic and clinical data of hMPV positive patients and hMPV negative patients.

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age group 4 to <5 years. Thus, the age and sex distribution of hMPV is described in Figure 4. Demographic and clinical characteristics of hMPV infections among children in this study are summarized in Table 4. Some significant differences were observed among demographic data. With the exception of few features, the clinical features showed no significant statistical differences between hMPV positive and hMPV negative patients.

Pneumonia and bronchiolitis were more frequent among hMPV positive patients than other group (P value <0.05).

The results of this study showed that the peak of hMPV infections was during February, which was 9 out of 48 (18.8%) hMPV positive patients, while the lowest frequency of infections was present in July which was negative for any hMPV infections. Therefore, autumn and winter is the highest season of hMPV infections, while summer is the least season of hMPV infections (Figure 5).

The calculation of sensitivity, specificity, and predictive values was performed for DFA depending on RT-PCR as a standard test. The results showed DFA sensitivity of 91.7%, specificity of 100%, positive predictive value of 100%, and negative predictive value of 98.4% (Table 5).

### Discussion

To the best of our knowledge, this is the first report for the detection of hMPV infection in Iraq using RT-PCR compared to Immunofluorescent technique. Seroprevalence of IgM and IgG anti-hMPV antibodies among hospitalized children under 5 years of age in Sulaimani city were 12% and 75%, respectively [5]. Consequently, human Metapneumovirus has shown to be responsible for a significant spectrum of clinical diseases with different manifestations in hMPV infections.

The general investigations showed that leukocytosis, subnormal SpO₂ %, and abnormal chest x-ray were present more frequently in hMPV positive children than in the other group. Hence, the differences between the two groups were statistically significant (Table 4). These findings show that hMPV infections are moderate to severe in potency.

Pneumonia and bronchiolitis were the main clinical diseases (70.8%) in hMPV positive children with 43.8% and 27%, respectively. Wei et al. reported bronchiolitis as the main clinical disease [19], while another study recorded pneumonia as the main clinical disease [23].

The study showed that hMPV infections were higher among age group <1 year (29.2%) when compared to older children. This result was comparable to that of Chen, et al. [24], Sohier Yahia, et al. [16], and Cui Zhang, et al. [25]. The higher incidence of hMPV infections among age group less than one year, might signify the importance of this age group as a risk factor for infection. However, Papenburg, et al. reported comparable results that children <6 months are at risk of hMPV infections [26].

In hMPV positive children, the male to female ratio was 2.4:1. Thus, this indicates male predominance, but the result was not significant when compared to hMPV negative children. Most published studies reported no significant gender predominance; Pilger, et al. recorded results similar to the current study [27], Zuo, et al. showed no sex difference among hMPV positive children [28], while Rein, et al. found that 53.2% of children infected with hMPV were females [29].

The results showed that 4 (13.3%) out of 30 of immunocompromized children were hMPV positive, while the remaining 26 (86.7%) were hMPV negative by DFA and RT-PCR. Therefore, the results were not statistically significant when compared to immunocompetent children. The clinical features of hMPV among the two groups showed no differences. Similar results were also obtained by John Williams et al., who found that patients infected with hMPV had varying degrees of immunosuppression, thereby suggesting that severe immune suppression is not a necessary risk factor for hMPV infection [30].

The results also clarified that living in rural regions was more in hMPV positive children than in the other group, and contact with respiratory infected patients was more frequent among hMPV positive children than in the second group. Comparable results were also found in other studies [31,32].

### Table 5: Sensitivity, specificity, and predictive values for the detection of hMPV in respiratory specimens.

<table>
<thead>
<tr>
<th>Parameter</th>
<th>DFA</th>
<th>RT-PCR</th>
</tr>
</thead>
<tbody>
<tr>
<td>True positive</td>
<td>44</td>
<td>48</td>
</tr>
<tr>
<td>False positive</td>
<td>0</td>
<td>0</td>
</tr>
<tr>
<td>True negative</td>
<td>252</td>
<td>252</td>
</tr>
<tr>
<td>False negative</td>
<td>4</td>
<td>0</td>
</tr>
<tr>
<td>Total number</td>
<td>300</td>
<td>300</td>
</tr>
<tr>
<td>Sensitivity</td>
<td>91.7%</td>
<td>100%</td>
</tr>
<tr>
<td>Specificity</td>
<td>100%</td>
<td>100%</td>
</tr>
<tr>
<td>Positive predictive value</td>
<td>100%</td>
<td>100%</td>
</tr>
<tr>
<td>Negative predictive value</td>
<td>98.4%</td>
<td>100%</td>
</tr>
</tbody>
</table>

![Figure 5: Seasonal distributions of hMPV infections among hospitalized children in the study group.](image)
The sample collection was done between April 2011 to March 2012, and the majority of HMMP positive patients were detected during the autumn and winter period with a peak in February. Other studies reported variable seasonal predominance. In a study done in China, spring was the main season [33]; in Jordan, the peak was reported in March [13]; in India, HMMP infection rate was at its peak in spring-summer period of 2008-2009 and 2009-2010; and HMMP circulated predominantly during the winter-spring period of 2010-2011 [34].

Conclusion

The HMMPV is an important cause of RTIs for hospitalized children in Sulaimani city, and both RT-PCR and DFA are valuable diagnostic techniques for this virus. However, RT-PCR is more sensitive than DFA. In addition, the autumn and winter period was the most frequent seasons for HMMPV infections.

Abbreviations

Human Metapneumovirus (hMPV), respiratory tract infections (RTIs), viral transport medium (VTM), Nasopharyngeal swabs (NPs), throat swabs (TSs), direct immunofluorescence assay (DFA), conventional reverse transcriptase – PCR (RT-PCR), Respiratory syncytial virus (RSV).

Acknowledgement

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References
